

(12) INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

(19) World Intellectual Property Organization
International Bureau(43) International Publication Date
8 November 2001 (08.11.2001)

PCT

(10) International Publication Number
WO 01/83774 A2

- (51) International Patent Classification⁷: C12N 15/55, 9/14, 5/10, A61K 38/46, C07H 1/00
- (21) International Application Number: PCT/US01/14116
- (22) International Filing Date: 2 May 2001 (02.05.2001)
- (25) Filing Language: English
- (26) Publication Language: English
- (30) Priority Data:
09/564,559 4 May 2000 (04.05.2000) US
- (71) Applicant (for all designated States except US):
PHOENIX PHARMACOLOGICS, INC. [US/US];
Suite A217, ASTeCC Facility, University of Kentucky,
Lexington, KY 40506-0286 (US).
- (72) Inventors; and
- (75) Inventors/Applicants (for US only): ENSOR, Charles, Mark [US/US]; 708 Dicksonia Court, Lexington, KY 40517 (US). HOLTSBERG, Frederick, Wayne [US/US]; 214 Garnet Drive, Nicholasville, KY 40356 (US). CLARK, Mike, A. [US/US]; 3940 Hall Road, Big Pine, FL 33043 (US).
- (74) Agents: ELDERKIN, Dianne, B. et al.; Woodcock Washburn Kurtz Mackiewicz & Norris LLP, 46th floor, One Liberty Place, Philadelphia, PA 19103 (US).
- (81) Designated States (national): AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, BZ, CA, CH, CN, CO, CR, CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, MZ, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW.
- (84) Designated States (regional): ARIPO patent (GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).
- Declaration under Rule 4.17:
— of inventorship (Rule 4.17(iv)) for US only
- Published:
— without international search report and to be republished upon receipt of that report
- For two-letter codes and other abbreviations, refer to the "Guidance Notes on Codes and Abbreviations" appearing at the beginning of each regular issue of the PCT Gazette.

(54) Title: NOVEL MUTATED FORM OF ARGININE DEIMINASE

Amino acid sequence of arginine deiminase (ADI) from wild-type *Mycoplasma hominis*

M. hominis ADI	MSVFDSKFNGLHVVYSEIGELETVLVHEPGREIDYITPARLDELLFSAILESHDARKEHQS	60
M. hominis ADI	FVKIMKDRGINVVELTDLVAETYDLASKAAKEEFIEFTLEETVPVLTEANKKAVRAFLLS	120
M. hominis ADI	KPTHEMVEFMMSGITKVELGVESENELJVDMPNLYFTRDPFASVGNVGTIHFMYTVRR	180
M. hominis ADI	RETLFARFVFRNHPKLKTPWYYDPAMKMPIEGGDVFIYNNETLVVGVSERTDLDTITLL	240
M. hominis ADI	AKNIKANKEVEFKRIVAINVPKWTNLMHLDTWLTMLDKNKFLYSPIANDVFKFWDYDLVN	300
M. hominis ADI	GGAEPQPQLNGLPLDKLLASINKEPVLPIGGAGATEMEIARETNFDGTNYLAIKPLGV	360
M. hominis ADI	IGYDRNEKTNAALKAAGITVLPFHGNQLSLGMGNARCSMPLSRKDVKW	409

(57) Abstract: The present invention discloses arginine deiminase that is genetically modified for more efficient manufacturing and processing. The present invention discloses recombinant DNA molecules and vectors and other therapeutic and pharmaceutical compositions. The present invention also discloses methods for preparing modified arginine deiminase as well as methods of treating cancer and other disease states using modified arginine deiminase.

WO 01/83774 A2

BEST AVAILABLE COPY

-1-

NOVEL MUTATED FORM OF ARGININE DEIMINASE

Field of the Invention

The present invention is directed to mutations of arginine deiminase for more efficient production and processing and thus improved treatment of cancer and other disease states.

Background of the Invention

It is presently believed that the amino acid arginine may play an important role in mediating certain disease states. For example, it has been reported that tumors such as melanomas, hepatomas, sarcomas and leukemia require arginine for growth. Sugimura et al., *Melanoma Res.* 2: 191-196 (1992); Takaku et al., *Int. J. Cancer* 51:244-249 (1992); Miyazaki et al., *Cancer Res.* 50:4522-4527 (1990); J.B. Jones, "The Effect of Arginine Deiminase on Murine Leukemic Lymphoblasts," Ph.D. Dissertation, The University of Oklahoma, pages 1-165 (1981). Malignant melanoma (stage 3) and hepatoma are fatal diseases which kill most patients within one year of diagnosis. In the United States, approximately 16,000 people die from these diseases annually. The incidence of melanoma is rapidly increasing in the United States and is even higher in other countries, such as Australia. The incidence of hepatoma, in parts of the world where hepatitis is endemic, is even greater. For example, hepatoma is one of the leading forms of cancer in Japan and Taiwan.

It has also been reported that many protozoa require arginine for growth and thus arginine may play an important role in many parasitic diseases. van Wagtenonk et al., "Nitrogen Metabolism in Protozoa", in *Comparative Biochemistry of Nitrogen Metabolism*, pages 1-56 (J.W. Campbell ed. 1970). Arginine (derived from the circulation) has also been shown to be a source of nitrous oxide which can play an important role in mediating septic shock. Chang et al., *Am. J. Physiol.* 274:H342-H348 (1998); McDonald et al., *J. Biol. Chem.* 272:31213-31216 (1997). Effective treatments for these diseases are urgently needed.

It has been reported that enzymes which degrade non-essential amino acids, such as arginine, may be an effective means of controlling some forms of cancer. For example, arginine deiminase isolated from *Pseudomonas putida* was described by J.B. Jones, "The Effect of Arginine Deiminase on Murine Leukemic Lymphoblasts," Ph.D. Dissertation, The University

-2-

of Oklahoma, pages 1-165 (1981). Because arginine deiminase catalyzes the conversion of arginine into citrulline, thus helping to eliminate arginine from the circulation of animals, it is believed that arginine deiminase can be used as an effective therapy for cancer and other disease states where arginine plays a role. Although arginine deiminase is not produced in mammals, it is found in a variety of bacteria, fungi and mycoplasma. Arginine deiminase can be thus be isolated from those organisms which produce it or, in the alternative, the enzyme may be produced using recombinant DNA technology. Misawa et al., *J. Biotechnol.* 36: 145-155 (1994).

Certain disadvantages have come to be associated with the isolation of arginine deiminase from organisms. Although effective in killing tumor cells *in vitro*, arginine deiminase isolated from *Pseudomonas putida* failed to exhibit efficacy *in vivo* because it had little enzyme activity at a neutral pH and was rapidly cleared from the circulation of experimental animals. Arginine deiminase derived from *Mycoplasma arginini* is described, for example, by Takaku et al, *Int. J. Cancer*, 51:244-249 (1992), and U.S. Patent No. 5,474,928, the disclosures of which are hereby incorporated by reference herein in their entirety. A problem associated with the therapeutic use of such a heterologous protein is its antigenicity. The chemical modification of arginine deiminase from *Mycoplasma arginini*, via a cyanuric chloride linking group, with polyethylene glycol was described by Takaku et al., *Int. J. Cancer Res.* 84:1195-1200 (1993). The modified protein was toxic when metabolized due to the release of cyanide from the cyanuric chloride linking group.

The production of arginine deiminase via recombinant DNA techniques also provides for certain disadvantages. For example, arginine deiminase produced in *Escheria coli* is enzymatically inactive and thus must be denatured and then properly renatured in order for it to become enzymatically active. The usual method for renaturing arginine deiminase produced in *E. coli* is to isolate the inactive enzyme, dissolve it in guanidium hydrochloride and renature it by rapid dilution into low ionic strength buffer. This last step requires very large volumes of buffer thus making the manufacture of arginine deiminase both expensive and time consuming. However, recombinant technology does have certain advantages. For example, organisms more amenable to fermentation can be used as hosts. Additionally, these fermentation hosts are generally much less pathogenic and larger amounts of arginine deiminase can be obtained. It has been shown the *E. coli* may produce large amounts of *Mycoplasma arginine deiminase*.

-3-

Another problem associated with arginine deiminase is that the enzyme is highly antigenic and thus rapidly cleared from circulation. Accordingly, arginine deiminase must be properly formulated before being used as a therapeutic agent. For purposes of the present invention, the term formulation can be defined as the chemical modification of any arginine deiminase for purposes of reducing antigenicity of the enzyme. For example, it has been shown that the formulation of several proteins including arginine deiminase with polyethylene glycol, i.e. pegylation, can reduce the antigenicity of the protein and greatly increase its circulating half-life. Unfortunately, the formulation of arginine deiminase with polyethylene glycol often inactivates the enzyme.

There is a need for methods and compounds which address these problems associated with the prior art. The present invention is directed to these, as well as other, important ends.

Summary of the Invention

The present invention relates to a modified arginine deiminase enzyme capable of more efficient production and processing.

The present invention further relates to a recombinant DNA molecule which encodes modified arginine deiminase.

In related aspects, the present invention further relates to recombinant vectors which comprise a nucleotide sequence that encodes modified arginine deiminase and to host cells comprising such vectors.

The present invention further relates to methods for preparing the modified arginine deiminase enzyme of the invention. These methods comprise growing a host cell transformed with the recombinant DNA molecule of this invention in a suitable culture medium.

In other aspects, this invention to methods of treating cancer as well as treating and/or inhibiting the metastasis of tumor cells. The invention also relates to methods of treating parasitic disease, septic shock and other disease states.

These and other aspects of the present invention will be elucidated in the following detailed description of the invention.

Brief Description of the Drawings

Figure 1 depicts the amino acid sequence of arginine deiminase cloned from wild-type *Mycoplasma hominus* (SEQ ID NO:1).

Figure 2 depicts the amino acid sequence of modified arginine deiminase from *Mycoplasma hominus* in accordance with preferred embodiments of the present invention (SEQ ID NO:4).

Detailed Description of Preferred Embodiments of the Invention

The present invention is based on the discovery that modifications of one or more of the naturally occurring amino acid residues of arginine deiminase from *Mycoplasma hominus* can provide for an enzyme that is more easily renatured and formulated thereby improving existing techniques for the manufacture of arginine deiminase and therapeutic compositions comprising the same. The therapeutic compositions of the invention may comprise recombinant DNA molecules, recombinant vectors including plasmids, transformed host cells and other pharmaceutical compositions. The therapeutic compositions may also comprise biocompatible carriers or diluents as known to those skilled in the art. The therapeutic compositions of the invention are easily made sterile and are non-pyrogenic. Such improved techniques and compositions are necessary for the effective treatment of cancer and other disease states.

As used herein, the term "melanoma" may be a malignant or benign tumor arising from the melanocytic system of the skin and other organs, including the oral cavity, esophagus, anal canal, vagina, leptomeninges, and/or the conjunctivae or eye. The term "melanoma" includes, for example, acral-lentiginous melanoma, amelanotic melanoma, benign juvenile melanoma, lentigo maligna melanoma, malignant melanoma, nodular melanoma, subungual melanoma and superficial spreading melanoma.

"Hepatoma" may be a malignant or benign tumor of the liver, including, for example, hepatocellular carcinoma.

"Patient" refers to an animal, preferably a mammal, more preferably a human.

-5-

"Biocompatible" refers to materials or compounds which are generally not injurious to biological functions and which will not result in any degree of unacceptable toxicity, including allergenic and disease states.

"Polyethylene glycol" or "PEG" refers to mixtures of condensation polymers of ethylene oxide and water, in a branched or straight chain, represented by the general formula $H(OCH_2CH_2)_nOH$, wherein n is at least 4. It is preferred that the polyethylene glycol is a straight chain. Generally, increasing the molecular weight of the polyethylene glycol decreases the immunogenicity of arginine deiminase. The polyethylene glycol may be used in conjunction with arginine deiminase, and, optionally, a biocompatible linking group, to treat cancer, including, for example, melanomas, hepatomas and sarcomas, preferably melanomas.

Normal cells do not require arginine for growth because they can synthesize arginine from citrulline in a two step process catalyzed by arginosuccinate synthase and arginosuccinate lyase. In contrast, melanomas, hepatomas and some sarcomas do not express arginosuccinate synthase; therefore, they are auxotrophic for arginine, i.e., they require arginine for growth. This metabolic difference may be capitalized upon to develop a safe and effective therapy to treat these forms of cancer. Arginine deiminase catalyzes the conversion of arginine to citrulline, and may be used to eliminate arginine. Thus, arginine deiminase may be utilized as a treatment for melanomas, hepatomas, some sarcomas and other disease states.

The amino acid sequences of arginine deiminase from the *Mycoplasma hominus* gene is disclosed by Figures 1 or 2. Chemical and genetic modification of the arginine deiminase enzyme can affect its biological activities. For example, it has been shown that arginine deiminase is typically antigenic and rapidly cleared from circulation in a patient. However, it has also been shown that the formulation of arginine deiminase with polyethylene glycol reduces the antigenicity and increases the circulating half-life of the enzyme. Abuchowski et al., *Cancer Biochem. Biophys.* 7:175-186 (1984); Abuchowski et al., *J. Biol. Chem.* 252:3582-3586 (1977). In particular, arginine deiminase can be covalently modified with polyethylene glycol. Arginine deiminase covalently modified with polyethylene glycol (with or without a linking group) may be hereinafter referred to as "ADI-PEG." In U.S. Patent Application Serial No. 09/023,809, Clark describes improved modifications of arginine deiminase with polyethylene glycol, the disclosure of which is hereby incorporated by reference herein in its entirety. When compared to native arginine deiminase, ADI-PEG retains most of its enzymatic

-6-

activity, is far less antigenic, has a greatly extended circulating half-life, and is much more efficacious in the treatment of tumors. For purposes of the invention, the modification of any arginine deiminase with polyethylene glycol may be referred to as pegylation.

It is necessary, however, to modify arginine deiminase with the maximum amount of polyethylene glycol in order to decrease antigenicity. For example, arginine deiminase can be formulated with the maximum amount of succinimidyl succinate polyethylene glycol (SS-PEG). The SS-PEG attaches to primary amines on proteins (the n-terminus and lysines) during the pegylation process. Unfortunately, it has been shown that the pegylation process can inactivate arginine deiminase if allowed to go to completion. While not meant to limit the present invention in any way, it is presently believed that arginine deiminase is inactivated because polyethylene glycol attaches to sites on the enzyme which interfere with the enzymes ability to catalyze a reaction. Thus, it has traditionally been necessary to carefully determine the exact ratio of SS-PEG to enzyme, concentration of reactant and time of reaction in order to prevent inactivation while decreasing antigenicity (i.e., optimal pegylation). Additionally, it becomes increasingly difficult to control pegylation reactions as manufacturing scale increases, and often it is necessary to inactivate the SS-PEG prior to removal of excess polyethylene glycol.

It has also been shown that arginine deiminase produced by recombinant technology, in particular arginine deiminase produced in *Escheria coli* cells, is initially inactive. It has thus been necessary to activate the recombinant arginine deiminase by denaturing and then properly renaturing the enzyme. Typically, the inactive enzyme has been isolated, dissolved in guanidium hydrochloride, and then renatured via rapid dilution with low ionic strength buffer. However, large volumes of buffer are required, thus making the production of recombinant arginine deiminase both expensive and time-consuming.

It has now been discovered that certain modifications of arginine deiminase can facilitate both the renaturation and formulation (i.e., pegylation) of the enzyme thereby improving manufacturing processes. The present invention is based on the unexpected discovery that certain amino acid changes in arginine deiminase provide for excellent results in the recombinant production and formulation of the enzyme. The production of arginine deiminase has traditionally been expensive and time consuming due to difficulties associated with recombinant production and formulation. The present invention thus discloses a modified

-7-

arginine deiminase that provides for improved renaturation and formulation processes while retaining the ability to convert arginine to citrulline. For purposes of the present invention, modified arginine deiminase can be defined as arginine deiminase having one or more amino acid substitutions. It is to be understood that modified arginine deiminase can include arginine deiminase having a single amino acid substitution or a plurality of amino acid substitutions. The invention also discloses a DNA sequence which encodes for the modified arginine deiminase of the invention. The modified arginine deiminase of the invention can provide for excellent results in the treatment of certain types of cancer, inhibiting the metastasis of cancer, and treating other disease states.

It is to be understood that preferred embodiments of the invention are based on the discovery that certain pegylation sites associated with arginine deiminase may be located at or adjacent the catalytic region of the enzyme. For purposes of the present invention, the phrase "pegylation site" may be defined as any site or position of arginine deiminase that may be covalently modified with polyethylene glycol. A "pegylation site" can be considered located at or adjacent the catalytic region of the enzyme where pegylation of the site results in a significant reduction in catalytic activity of the enzyme. The pegylation of such sites has traditionally resulted in the inactivation of the enzyme. For example, arginine deiminase from *Mycoplasma hominus* has a lysine at the 112 position which is believed to be at or adjacent the catalytic region of the enzyme. The attachment of polyethylene glycol to this lysine at the 112 position may inactivate the enzyme. In addition, arginine deiminase from *Mycoplasma hominus* has a cysteine at the 397 position which is believed to be at or adjacent the catalytic region of the enzyme. The importance of this cysteine has been shown as it has now been discovered that amino acid substitutions for cysteine at the 397 position may inactivate the enzyme. In particular, it has been shown that substituting alanine, histidine, arginine, serine, lysine or tyrosine for cysteine at the 397 position results in a loss of all detectable enzyme activity. Arginine deiminase from *Mycoplasma hominus* also has three lysines located near this conserved cysteine, in particular Lys³⁷⁴, Lys⁴⁰⁵ and Lys⁴⁰⁸. The attachment of polyethylene glycol to Lys³⁷⁴, Lys⁴⁰⁵, Lys⁴⁰⁸ or combinations thereof is believed to inactivate the enzyme.

It is to be understood that arginine deiminase derived from other organisms may also have pegylation sites corresponding to 112 position of arginine deiminase from *Mycoplasma hominus*. For example, arginine deiminase from *Streptococcus pyrogenes* has lysine

at the 104 position, arginine deiminase from *Mycoplasma pneumoniae* has lysine at the 106 position, and arginine deiminase from *Giardia intestinalis* has lysine at the 114 position. In addition, arginine deiminase from some organisms may have lysines corresponding to the same general location as the 112 position of arginine deiminase from *Mycoplasma hominus*. The location of lysine in arginine deiminase from such organisms may be indicated as follows:

Table 1: Pegylation sites of arginine deiminase from various organisms

Organisms producing arginine deiminase	Position of lysine in arginine deiminase
<i>Mycoplasma hominus</i>	112
<i>Mycoplasma arginini</i>	111
<i>Clostridium perfringens</i>	105
<i>Bacillus licheniformis</i>	97, 108
<i>Borrelia burgdorferi</i>	102, 111
<i>Borrelia afzelii</i>	101
<i>Enterococcus faecalis</i>	102, 110
<i>Streptococcus pyogenes</i>	104
<i>Streptococcus pneumoniae</i>	103
<i>Lactobacillus sake</i>	97, 106
<i>Giardia intestinalis</i>	114, 116

It is presently believed that the attachment of polyethylene glycol to such lysines or combinations thereof may inactivate the enzyme. It is presently believed that amino acid substitutions at such lysines may result in a protein that loses less of its enzymatic activity upon pegylation.

The present invention thus provides for certain amino acid substitutions in the polypeptide chain of arginine deiminase. These amino acid substitutions provide for modified arginine deiminase that loses less activity upon pegylation; i.e. upon pegylation, the reduction of enzyme activity following pegylation in the modified arginine deiminases is less than the reduction of enzyme activity following pegylation in the unmodified arginine deiminases. By eliminating pegylation sites at or adjacent to the catalytic region of enzyme,

-9-

optimal pegylation can be achieved without the traditional loss of activity. As discussed above, arginine deiminase from certain organisms have pegylation sites located at various positions on the peptide chain. While not limiting the present invention, it is presently believed that arginine deiminase may have the amino acid lysine located at or adjacent to the catalytic region of the enzyme and that pegylation of these sites may inactivate the enzyme. By eliminating at least one of these pegylation sites, pegylation can be achieved and more enzyme activity retained. In accordance with the invention, it is preferred that lysine is substituted with glutamic acid, valine, aspartic acid, alanine, isoleucine, leucine or combinations thereof. More preferred is that lysine is substituted with glutamic acid. In one embodiment of the invention, it is preferred that modified arginine deiminase from *Mycoplasma hominus* has an amino acid substitution at Lys¹¹² (SEQ ID NO:2), Lys³⁷⁴, Lys⁴⁰⁵, Lys⁴⁰⁸ or combinations thereof. Preferably, modified arginine deiminase from *Mycoplasma hominus* has an amino acid substitution Lys¹¹² to Glu¹¹², Lys³⁷⁴ to Glu³⁷⁴, Lys⁴⁰⁵ to Glu⁴⁰⁵, Lys⁴⁰⁸ to Glu⁴⁰⁸ or combinations thereof. More preferred is that modified arginine deiminase from *Mycoplasma hominus* has lysine at position 112 substituted with glutamic acid.

It is to be understood that other preferred embodiments of the invention are based on the discovery that certain structural characteristics of arginine deiminase may prevent or interfere with the proper and rapid renaturation of arginine deiminase when produced via recombinant technology. In particular, these structural characteristics hinder or prevent the enzyme from assuming an active conformation during recombinant production. For purposes of the present invention, the phrase "active conformation" may be defined as a three-dimensional structure that allows for enzymatic activity by unmodified or modified arginine deiminase. The active conformation may, in particular, be necessary for catalyzing the conversion of arginine into citrulline. The phrase "structural characteristic" may be defined as any trait, quality or property of the polypeptide chain resulting from a particular amino acid or combination of amino acids. For instance, arginine deiminase may contain an amino acid that results in a bend or kink in the normal peptide chain and thus hinders the enzyme from assuming an active conformation during renaturation of the enzyme. In particular, arginine deiminase from *Mycoplasma hominus* has a proline at the 210 position that may result in a bend or kink in the peptide chain, making it more difficult to renature the enzyme during recombinant production. It is to be understood

-10-

that arginine deiminase derived from other organisms may also have sites corresponding to the 210 position of arginine deiminase from *Mycoplasma hominus*.

The present invention thus again provides for certain amino acid substitutions in the polypeptide chain of arginine deiminase. Such amino acid substitutions can eliminate the problematic structural characteristics in the peptide chain of arginine deiminase. Such amino acid substitutions provide for improved renaturation of the modified arginine deiminase. These amino acid substitutions make possible rapid renaturing of modified arginine deiminase using reduced amounts of buffer. These amino acid substitutions may also provide for increased yields of renatured modified arginine deiminase. In one embodiment of the invention, it is preferred that the modified arginine deiminase have a single amino acid substitution at Pro²¹⁰. As mentioned above, arginine deiminase derived from *Mycoplasma hominus* has the amino acid proline located at the 210 position. While not limiting the present invention, it is presently believed that the presence of the amino acid proline at position 210 results in a bend or kink in the normal polypeptide chain that increases the difficulty of renaturing (i.e., refolding) arginine deiminase. Substitutions for proline at position 210 make possible the rapid renaturation of modified arginine deiminase using reduced amounts of buffer. Substitutions for proline at position 210 may also provide for increased yields of renatured modified arginine deiminase. In a preferred embodiment, the proline at position 210 is substituted with serine (SEQ ID NO:3). It is to be understood that in accordance with this aspect of the invention, other substitutions at position 210 may be made. Examples of preferred substitutions include Pro²¹⁰ to Thr²¹⁰, Pro²¹⁰ to Arg²¹⁰, Pro²¹⁰ to Asn²¹⁰, Pro²¹⁰ to Gln²¹⁰ or Pro²¹⁰ to Met²¹⁰. By eliminating those structural characteristics associated with the amino acid of position 210 of the wild-type arginine deiminase, proper refolding of the enzyme can be achieved.

In another embodiment of the invention, it is preferred that the modified arginine deiminase have multiple amino acid substitutions. The modified arginine deiminase may have at least one amino acid substitution eliminating pegylation sites at or adjacent a catalytic region of the enzyme. The modified arginine deiminase may also have at least one amino acid substitution eliminating those structural characteristics that interfere with the renaturation of the enzyme. The amino acid substitutions may thus provide for a modified arginine deiminase of the invention. The amino acid substitutions may provide for the pegylation of modified arginine deiminase without a loss of enzymatic activity. The amino acid

-11-

substitutions may provide for a modified arginine deiminase that can be rapidly renatured using reduced amounts of buffer. The amino acid substitutions may also provide for increased yields of renatured modified arginine deiminase. In a preferred embodiment, the modified arginine deiminase derived from *Mycoplasma hominus* includes the proline at position 210 substituted with serine and the lysine at position 112 substituted with glutamic acid (SEQ ID NO:4). As discussed above, however, it is to be understood that the modified arginine deiminase may include other preferred substitutions.

It is preferred that the modified arginine deiminase of the invention is derived from *Mycoplasma hominus*. In accordance with the invention, however, it will be understood by those skilled in the art that modified arginine deiminase may be derived from other organisms. For example, as discussed above, modified arginine deiminase of the invention can be derived from *Streptococcus pyogenes*, *Mycoplasma pneumoniae*, *Giardia intestinalis*, *Mycoplasma hominus*, *Mycoplasma arginini*, *Clostridium perfringens*, *Bacillus licheniformis*, *Borrelia burgdorferi*, *Borrelia afzelii*, *Enterococcus faecalis*, *Streptococcus pyogenes*, *Streptococcus pneumoniae*, *Lactobacillus sake*, *Giardia intestinalis*. It is to be understood that modified arginine deiminase may have one or more amino acid substitutions, in accordance with the various aspects of the invention.

It is to be understood that in certain embodiments of the present invention, modified arginine deiminase can be formulated with polyethylene glycol. The present invention provides for the attachment of polyethylene glycol to modified arginine deiminase for increasing circulation half-life, without the traditional inactivation of the enzyme. Traditionally, selection of the attachment site of polyethylene glycol on arginine deiminase was determined by the role of each of the sites within the active domain of the protein, as would be known to the skilled artisan. In accordance with the present invention, polyethylene glycol may be attached to a primary amine of the modified arginine deiminase. The present invention provides for the attachment of polyethylene glycol to the primary amines of arginine deiminase without substantial loss of enzymatic activity. For example, modified arginine deiminase from *Mycoplasma hominus* contains lysine amino acids that may be modified with polyethylene glycol. In other words, these lysines are all possible points at which modified arginine deiminase can be attached to polyethylene glycol. It is to be understood, however, that polyethylene glycol may also be attached to other sites on modified arginine deiminase, as would be apparent to those

-12-

skilled in the art. Increasing the number of polyethylene glycol units on modified arginine deiminase increases the circulating half life of the enzyme without the traditional decrease in specific activity of the enzyme.

The linking group used to covalently attach polyethylene glycol to modified arginine deiminase may be any biocompatible linking group. As discussed above, "biocompatible" indicates that the compound or group is non-toxic and may be utilized *in vitro* or *in vivo* without causing injury, sickness, disease or death. Polyethylene glycol can be bonded to the linking group, for example, via an ether bond, an ester bond, a thiol bond or an amide bond. Suitable biocompatible linking groups include, for example, an ester group, an amide group, an imide group, a carbamate group, a carboxyl group, a hydroxyl group, a carbohydrate, a maleimide group (including, for example, succinimidyl succinate (SS), succinimidyl propionate (SPA), succinimidyl carboxymethylate (SCM), succinimidyl succinamide (SSA) or N-hydroxy succinimide (NHS)), an epoxide group, an oxycarbonylimidazole group (including, for example, carbonyldimidazole (CDI)), a nitro phenyl group (including, for example, nitrophenyl carbonate (NPC) or trichlorophenyl carbonate (TPC)), a trysylate group, an aldehyde group, an isocyanate group, a vinylsulfone group, a tyrosine group, a cysteine group, a histidine group or a primary amine. Preferably, the biocompatible linking group is an ester group and/or a maleimide group. More preferably, the linking group is SS, SPA, SCM, SSA or NHS; with SS, SPA or NHS being more preferred, and with SS or SPA being most preferred.

Alternatively, polyethylene glycol may be coupled directly to modified arginine deiminase (i.e., without a linking group) through an amino group, a sulphydryl group, a hydroxyl group or a carboxyl group. Polyethylene glycol may be covalently bonded to modified arginine deiminase via a biocompatible linking group, using methods known in the art, as described, for example, by Park et al, *Cancer Res.*, 33:3-14 (1973); and Zaplipsky and Lee, *Polyethylene Glycol Chemistry: Biotechnical and Biomedical Applications*, J.M. Harris, ed., Plenum Press, NY, Chapter 21 (1992), the disclosures of which are hereby incorporated by reference herein in their entirety.

The present invention also relates to recombinant DNA molecules encoding the modified arginine deiminase of the invention. It is preferred that the DNA molecules comprising the recombinant DNA molecules of this invention are derived from any *Mycoplasma hominus* strain using known techniques, e.g., isolating the gene from a gene bank, making

-13-

complementary or cDNAs from a mRNA template or via the polymerase chain reaction (see, U.S. Patent No. 4,800,159) or from isolates of clinical specimens. Alternatively, such recombinant DNA molecules may be synthesized by standard DNA synthesis techniques. Various *Mycoplasma hominus* strains are also publicly available from commercial depositories, e.g., from the American Type Culture Collection (ATCC), Bethesda, MD, U.S.A.

The DNA molecules of this invention may comprise additional DNA sequences including, for example, a regulatory element, one or more selectable markers, and sequences that code for replication and maintenance functions. The regulatory region typically contains a promoter found upstream from the coding sequence of this invention, which functions in the binding of RNA polymerase and in the initiation of RNA transcription. In other words, the regulatory element or region can be operatively linked to the coding sequence of this invention. It will be appreciated by one of skill in the art that the selection of regulatory regions will depend upon the host cell employed.

The invention also relates to recombinant vectors, particularly recombinant plasmids that comprise sequences encoding the modified arginine deiminase of the invention.

Another aspect of this invention is a host cell transformed with the recombinant DNA molecule of this invention. Such host cell is capable of growth in a suitable culture medium and expressing the coding sequence of the invention. Such host cell can be prepared by methods of this invention, e.g., by transforming a desired host cell with the plasmid of this invention. Such transformation can be accomplished by utilization of conventional transformation techniques. Furthermore, the recombinant DNA molecule of this invention can be integrated into the host cell's genome by conventional techniques, e.g., homologous recombination. Those host cells suitable for use in the present invention include, but are not limited to, mammalian cells, insect cells, yeast and other bacteria cells, e.g., *Streptomyces*, *Bacillus* and *Salmonella*. It is preferred that the host cells of this invention include those belonging to the species *E. coli*. This invention and the product thereof is not limited to any specific host cell. It is preferred that the modified arginine deiminase is produced by the transformed host cell of this invention, but such enzyme can be prepared by conventional peptide synthesis techniques.

The present invention also relates to methods of producing the enzyme encoded by the recombinant DNA molecule of this invention which comprises culturing the

-14-

transformed host of the invention in an appropriate culture media and the isolation of such enzyme. For purposes of the present invention, the phrase "appropriate culture media" can be defined as that media which facilitates such host in expressing recombinant DNA molecules encoding the modified arginine deiminase of the invention. It will be appreciated by those skilled in the art that the appropriate culture media will depend upon the host cell used. The isolation of the enzyme so produced can be accomplished from a culture lysate of the host, or if appropriate, directly from the host's culture medium, and such isolation is carried out by conventional protein isolation techniques.

In a preferred embodiment, the DNA coding sequence of the modified arginine deiminase of the invention is expressed in a transformed *Mycoplasma hominus* host cell. Preferably the *Mycoplasma hominus* cell is deficient in arginine deiminase and thus requires complementation. In such systems, sequences that encode the modified arginine deiminase are typically located on a vector. Such vectors contain sufficient amount of bacterial DNA to propagate the vector in *E. coli* or some other suitable host. Such vector also contains a sufficient amount of *Mycoplasma hominus* DNA flanking the enzyme coding sequence so as to permit recombination between a *Mycoplasma hominus* host deficient in the arginine deiminase gene and the heterologous modified arginine deiminase gene. It is to be understood by those skilled in the art that it is not essential to use a *Mycoplasma hominus* host deficient in the modified arginine deiminase gene, but that the absence of the gene in the host prior to recombination will facilitate the screening and isolation of recombinant hosts which have incorporated the gene of interest. The recombinant *Mycoplasma hominus* arising from such homologous recombination can then be selected by standard techniques as known to those skilled in the art.

The invention also encompasses methods for the treatment of cancer and/or other disease states comprising the administration of a therapeutically effective amount of one of the compounds of the present invention. A therapeutically effective amount of one of the compounds of the present invention is an amount that is effective in reducing the incidence of the disease state. Where the disease state is cancer, a therapeutically effective amount of one of the compounds of the present invention can be an amount that is effective to inhibit tumor growth. Generally, treatment is initiated with small dosages which can be increased by small increments until the optimum effect under the circumstances is achieved. Generally, a therapeutic dosage of compounds of the present invention may be from about 1 to about 200

-15-

mg/kg twice a week to about once every two weeks. For example, the dosage may be about 1 mg/kg once a week as a 2 ml intravenous injection to about 20 mg/kg once every 3 days. PEG-ADI may be mixed with a phosphate buffered saline solution, or any other appropriate solution known to those skilled in the art, prior to injection. The PEG-ADI formulation may be administered as a solid (lyophilate) or as a liquid formulation, as desired.

The methods of the present invention can involve either *in vitro* or *in vivo* applications. In the case of *in vitro* applications, including cell culture applications, the compounds described herein can be added to the cells in cultures and then incubated. The compounds of the present invention may also be used to facilitate the production of monoclonal and/or polyclonal antibodies, using antibody production techniques well known in the art. The monoclonal and/or polyclonal antibodies can then be used in a wide variety of diagnostic applications, as would be apparent to one skilled in the art.

The *in vivo* means of administration of the compounds of the present invention will vary depending upon the intended application. As one skilled in the art will recognize, administration of the PEG-ADI composition of the present invention can be carried out, for example, orally, intranasally, intraperitoneally, parenterally, intravenously, intralymphatically, intratumorally, intramuscularly, interstitially, intra-arterially, subcutaneously, intraocularly, intrasynovial, transepithelial, and transdermally.

Examples

The invention is further demonstrated in the following examples, which are for purposes of illustration, and are not intended to limit the scope of the present invention.

Example 1: *Cloning and Site-directed mutagenesis of ADI*

Cultures of *Mycoplasma hominus* (ATCC 23114) were obtained from the American Type Culture, Bethesda, MD. DNA was extracted and the gene coding for arginine deiminase isolated by the polymerase chain reaction.

The arginine deiminase gene was subcloned into pGEM-t and the *E. coli* strain JM101. Site directed mutagenesis was performed using the Altered Sites II kit (Promega, Madison, Wisconsin).

-16-

Modified arginine deiminase was expressed in JM101 cells as previously described by Takaku et al., *supra*. The modified arginine deiminase included glutamic acid at the 112 position and serine at the 210 position (SEQ ID NO:4). The amino acid sequence of modified arginine deiminase from *Mycoplasma hominus* is described in Figure 2.

Example 2: Renaturation and purification of enzymatically active ADI

The modified arginine deiminase was isolated and purified as previously described by Takaku et al., *supra*. However, several improvements were observed.

Table 2: Renaturation of Arginine Deiminase

Compound	Time (° C)	Dilution Ratio	Yield
Arginine Deiminase	90 hours (15 ° C)	1:200	70 mg/L
Modified Arginine Deiminase	6-12 hours (room temp.)	1:50	500 mg/L

As indicated by Table 2 above, renaturation of the modified arginine deiminase was completed at room temperature in about 6 to 12 hours using a 1:50 dilution ratio of guanidium hydrochloride inclusion bodies in buffer. In contrast, Takaku et al. reported that renaturation required 90 hours at 15 ° C using a 1:200 dilution ratio. In addition, the yield of modified arginine deiminase was routinely about 500 mg per liter of fermentation whereas Takaku et al. reported a yields of approximately 70 mg per liter of fermentation.

Example 3: Formulation of modified ADI with PEG

Modified arginine deiminase was formulated using SS-PEG as previously described. The pegylation process was allowed to go unchecked for over 4 hours without the modified arginine deiminase becoming inactivated. In addition, it was not necessary to quench the pegylation process through the addition of glycine. With reference to Table 3, approximately 70-80% of enzymatic activity of the modified arginine deiminase was retained.

Table 3: Specific Enzyme Activity (IU/mg protein)

	Without Pegylation	Pegylated
wild type ADI	20-21	5-8
modified ADI	20-21	12-16

This indicates that the modified arginine deiminase allows for more consistent formulations. In addition, the modified arginine deiminase allows for scaling up of the manufacturing process as compared to the wild-type arginine deiminase.

Each of the patents, patent applications and publications described herein are hereby incorporated by reference in their entirety.

Various modifications of the invention, in addition to those described herein, will be apparent to one skilled in the art in view of the foregoing description. Such modifications are also intended to fall within the scope of the appended claims.

It is to be understood that the specification of the present application hereby incorporates by reference the claims and their disclosures in their entirety.

-18-

What is claimed is:

1. A modified arginine deiminase for improved manufacturing processes comprising arginine deiminase modified to be free of at least one pegylation site at or adjacent to its catalytic region.
2. The modified arginine deiminase of claim 1 wherein the arginine deiminase is derived from *Mycoplasma hominus*.
3. The modified arginine deiminase of claim 2 wherein the modification comprises the deletion or substitution of at least one amino acid.
4. The modified arginine deiminase of claim 3 wherein the modification comprises at least one amino acid substitution at the 112, 374, 405 or 408 positions.
5. The modified arginine deiminase of claim 4 wherein the modification comprises at least one amino acid substitution selected from the group comprising Lys¹¹² to Glu¹¹², Lys¹¹² to Val¹¹², Lys¹¹² to Asp¹¹², Lys¹¹² to Ala¹¹², Lys¹¹² to Ile¹¹², Lys¹¹² to Leu¹¹², Lys³⁷⁴ to Glu³⁷⁴, Lys³⁷⁴ to Val³⁷⁴, Lys³⁷⁴ to Asp³⁷⁴, Lys³⁷⁴ to Ala³⁷⁴, Lys³⁷⁴ to Ile³⁷⁴, Lys³⁷⁴ to Leu³⁷⁴, Lys⁴⁰⁵ to Glu⁴⁰⁵, Lys⁴⁰⁵ to Val⁴⁰⁵, Lys⁴⁰⁵ to Asp⁴⁰⁵, Lys⁴⁰⁵ to Ala⁴⁰⁵, Lys⁴⁰⁵ to Ile⁴⁰⁵, Lys⁴⁰⁵ to Leu⁴⁰⁵, Lys⁴⁰⁸ to Glu⁴⁰⁸, Lys⁴⁰⁸ to Val⁴⁰⁸, Lys⁴⁰⁸ to Asp⁴⁰⁸, Lys⁴⁰⁸ to Ala⁴⁰⁸, Lys⁴⁰⁸ to Ile⁴⁰⁸, Lys⁴⁰⁸ to Leu⁴⁰⁸, and combinations thereof.
6. The modified arginine deiminase of claim 2 wherein the modification comprises the substitution of lysine at the 112 position.
7. The modified arginine deiminase of claim 6 wherein the modification comprises an amino acid substitution selected from a group comprising Lys¹¹² to Glu¹¹², Lys¹¹² to Val¹¹², Lys¹¹² to Asp¹¹², Lys¹¹² to Ala¹¹², Lys¹¹² to Ile¹¹² and Lys¹¹² to Leu¹¹².

-19-

8. The modified arginine deiminase of claim 6 wherein the modification comprises the amino acid substitution Lys¹¹² to Glu¹¹².
9. The modified arginine deiminase of claim 1 wherein the arginine deiminase is derived from an organism selected from a group comprising *Mycoplasma hominus*, *Mycoplasma pneumoniae*, *Mycoplasma arginini*, *Qiardia intestinalis*, *Clostridium perfringens*, *Bacillus licheniformis*, *Borrelia burgdorferi*, *Borrelia afzellii*, *Enterococcus faecalis*, *Streptococcus pyogenes*, *Streptococcus pneumoniae*, *Lactobacillus sake* and *Qiardia intestinalis*.
10. The modified arginine deiminase of claim 9 wherein the modification comprises the deletion or substitution of at least one amino acid.
11. The modified arginine deiminase of claim 2 further comprising the arginine deiminase having a substitution of proline at the 210 position.
12. The modified arginine deiminase of claim 11 wherein the substitution of proline at the 210 position is selected from the group comprising Pro²¹⁰ to Ser²¹⁰, Pro²¹⁰ to Thr²¹⁰, Pro²¹⁰ to Arg²¹⁰, Pro²¹⁰ to Asn²¹⁰, Pro²¹⁰ to Gln²¹⁰ and Pro²¹⁰ to Met²¹⁰.
13. The modified arginine deiminase of claim 11 wherein the substitution of proline at the 210 position is Pro²¹⁰ to Ser²¹⁰.
14. The modified arginine deiminase of claim 11 wherein the modification comprises the deletion or substitution of at least one amino acid.

-20-

15. The modified arginine deiminase of claim 14 wherein the modification comprises at least one amino acid substitution at the 112, 374, 405 or 408 positions.
16. The modified arginine deiminase of claim 15 wherein the modification comprises at least one amino acid substitution selected from the group comprising Lys¹¹² to Glu¹¹², Lys¹¹² to Val¹¹², Lys¹¹² to Asp¹¹², Lys¹¹² to Ala¹¹², Lys¹¹² to Ile¹¹², Lys¹¹² to Leu¹¹², Lys³⁷⁴ to Glu³⁷⁴, Lys³⁷⁴ to Val³⁷⁴, Lys³⁷⁴ to Asp³⁷⁴, Lys³⁷⁴ to Ala³⁷⁴, Lys³⁷⁴ to Ile³⁷⁴, Lys³⁷⁴ to Leu³⁷⁴, Lys⁴⁰⁵ to Glu⁴⁰⁵, Lys⁴⁰⁵ to Val⁴⁰⁵, Lys⁴⁰⁵ to Asp⁴⁰⁵, Lys⁴⁰⁵ to Ala⁴⁰⁵, Lys⁴⁰⁵ to Ile⁴⁰⁵, Lys⁴⁰⁵ to Leu⁴⁰⁵, Lys⁴⁰⁸ to Glu⁴⁰⁸, Lys⁴⁰⁸ to Val⁴⁰⁸, Lys⁴⁰⁸ to Asp⁴⁰⁸, Lys⁴⁰⁸ to Ala⁴⁰⁸, Lys⁴⁰⁸ to Ile⁴⁰⁸, Lys⁴⁰⁸ to Leu⁴⁰⁸, and combinations thereof.
17. The modified arginine deiminase of claim 11 wherein the modification comprises the substitution of lysine at the 112 position.
18. The modified arginine deiminase of claim 17 wherein the modification comprises an amino acid substitution selected from a group comprising Lys¹¹² to Glu¹¹², Lys¹¹² to Val¹¹², Lys¹¹² to Asp¹¹², Lys¹¹² to Ala¹¹², Lys¹¹² to Ile¹¹² and Lys¹¹² to Leu¹¹².
19. The modified arginine deiminase of claim 17 wherein the modification comprises the amino acid substitution Lys¹¹² to Glu¹¹².
20. An isolated DNA molecule comprising a recombinant coding sequence encoding the modified arginine deiminase of claim 1.
21. A transformed host cell comprising the DNA molecule of claim 20.
22. A recombinant plasmid comprising the DNA molecule of claim 20.

-21-

23. A host cell transformed with the recombinant plasmid of claim 22.
24. A composition comprising the modified arginine deiminase of claim 1 covalently bound to polyethylene glycol
25. The composition of claim 24 wherein the modified arginine deiminase is covalently bound via a linking group to polyethylene glycol.
26. A pharmaceutical composition comprising the modified arginine deiminase of claim 24 and a pharmaceutically acceptable carrier or diluent.
27. A method for preparing modified arginine deiminase, comprising:
- (a) transforming a host cell with the DNA of claim 20;
 - (b) growing the transformed host cell in a suitable culture medium;
 - (c) isolating the modified arginine deiminase.
28. The method of claim 27 further comprising covalently bonding polyethylene glycol to the arginine deiminase.
29. An isolated DNA molecule comprising a recombinant coding sequence encoding the modified arginine deiminase of claim 11.
30. A transformed host cell comprising the DNA molecule of claim 29.
31. A recombinant plasmid comprising the DNA molecule of claim 29.
32. A host cell transformed with the recombinant plasmid of claim 31.

-22-

33. A composition comprising the modified arginine deiminase of claim 11 covalently bound to polyethylene glycol.
34. The composition of claim 33 wherein the modified arginine deiminase is covalently bound via a linking group to polyethylene glycol.
35. A pharmaceutical composition comprising the modified arginine deiminase of claim 33 and a biocompatible carrier or diluent.
36. A method for preparing modified arginine deiminase, comprising:
- (a) transforming a host cell with the DNA of claim 29;
 - (b) growing the transformed host cell in a suitable culture medium;
 - (c) isolating the modified arginine deiminase.
37. The method of claim 36 further comprising covalently bonding polyethylene glycol to the arginine deiminase.
38. A modified arginine deiminase from *Mycoplasma hominus* having a substitution of lysine to glutamic acid at the 112 position and a substitution of proline to serine at the 210 position.
39. An isolated DNA molecule comprising a recombinant coding sequence encoding the modified arginine deiminase of claim 38.
40. A transformed host cell comprising the DNA molecule of claim 39.
41. A recombinant plasmid comprising the DNA molecule of claim 39.
42. A host cell transformed with the recombinant plasmid of claim 41.

-23-

43. A composition comprising the modified arginine deiminase of claim 38 covalently bound to polyethylene glycol
44. The composition of claim 43 wherein the modified arginine deiminase is covalently bound via a linking group to polyethylene glycol.
45. A pharmaceutical composition comprising the modified arginine deiminase of claim 38 and a pharmaceutically acceptable carrier or diluent.
46. A method for preparing modified arginine deiminase, comprising:
- (a) transforming a host cell with the DNA of claim 39;
 - (b) growing the transformed host cell in a suitable culture medium;
 - (c) isolating the modified arginine deiminase.
47. The method of claim 46 further comprising covalently bonding polyethylene glycol to the modified arginine deiminase.
48. An isolated DNA molecule comprising a recombinant coding sequence encoding the modified arginine deiminase of claim 7.
49. A transformed host cell comprising the DNA molecule of claim 48.
50. A recombinant plasmid comprising the DNA molecule of claim 48.
51. A host cell transformed with the recombinant plasmid of claim 50.
52. A composition comprising the modified arginine deiminase of claim 48 covalently bound to polyethylene glycol.

-24-

53. The composition of claim 52 wherein the modified arginine deiminase is covalently bound via a linking group to polyethylene glycol.
54. A pharmaceutical composition comprising the modified arginine deiminase of claim 52 and a pharmaceutically acceptable carrier or diluent.
55. A method for preparing the modified arginine deiminase comprising:
- (a) transforming a host cell with the DNA molecule of claim 48;
 - (b) growing said transformed host cell in a suitable culture medium;
 - (c) isolating the modified arginine deiminase.
56. The method of claim 55 further comprising covalently bonding polyethylene glycol to the modified arginine deiminase.
57. A modified arginine deiminase from *Mycoplasma hominus* having a substitution of proline to serine at the 210 position.
58. An isolated DNA molecule comprising a recombinant coding sequence encoding the modified arginine deiminase of claim 57.
59. A transformed host cell comprising the DNA molecule of claim 58.
60. A recombinant plasmid comprising the DNA molecule of claim 58.
61. A host cell transformed with the recombinant plasmid of claim 60.
62. A composition comprising the modified arginine deiminase of claim 57 covalently bound to polyethylene glycol.
63. The compositions of claim 62 wherein the modified arginine deiminase is covalently bound via a linking group to polyethylene glycol

-25-

64. A pharmaceutical composition comprising the modified arginine deiminase of claim 57 and a pharmaceutically acceptable carrier or diluent.
65. A method for preparing the modified arginine deiminase comprising:
- (a) transforming a host cell with the DNA molecule of claim 58;
 - (b) growing said transformed host cell in a suitable culture medium;
 - (c) isolating the modified arginine deiminase.
66. The method of claim 65 further comprising covalently bonding polyethylene glycol to the modified arginine deiminase.
67. A method of treating a tumor in a patient comprising administering to said patient the composition of claims 24, 26, 33, 35, 43, 45, 52, 54, 62 or 64.
68. A method of treating and inhibiting metastases in a patient comprising administering to said patient the composition of claims 24, 26, 33, 35, 43, 45, 52, 54, 62 or 64.
69. A method of treating parasitic disease in a patient comprising administering to said patient the composition of claims 24, 26, 33, 35, 43, 45, 52, 54, 62 or 64.
70. A method of treating septic shock in a patient comprising administering to said patient the composition of claims 24, 26, 33, 35, 43, 45, 52, 54, 62 or 64.

FIG 1. Amino acid sequence of arginine deiminase (ADI) from wild-type *Mycoplasma hominus*

M. hominis ADI	MSVFDSKFNHVVYSEIGELETVLVHEPGREIDYITPARLDELLFSAILESHDARKEHQ	60
M. hominis ADI	FVKIMKDRGINVVELTDLVAETYDLASKAAKEEFIEFLEETVPVLTEANKKAVRAFLS	120
M. hominis ADI	KPTHEMVEFMMSGITKYELGVESENELIVDPMPNLYFTRDPFASVGNGVTHFMRYIVRR	180
M. hominis ADI	RETLFARFVFRNHPKLVKTPWYYDPAMKMPIEGDVFYNNETLVGVSEKTDLDITILL	240
M. hominis ADI	AKNIKANKEVEFKRIVAINVPKWTNLMHLDTWLTMLDKNKFLYSPIANDVFKFWDYDLVN	300
M. hominis ADI	GGAEPQQLNGLPLDKLLASINKEPVLPIGGAGATEMEIARETNFDGTNYLAKPGLV	360
M. hominis ADI	IGYDRNEKTNAALKAAGITVLPFHGNQLSLGMGNARCMSMPLSRKDVKW	409

FIG 2. Amino acid sequence of modified arginine deiminase (ADI E112, S210) from *Mycoplasma hominus*

ADI E112, S210	MSVFDSKFNHGVYSEIGELETVLVHEPGREIDYTPARLDELLFSAILSHDARKEHQS	60
ADI E112, S210	FVKIMKDRGINVVELTDLVAETYDLASKAAKEEFIEETVPVLTEANKEAVRAFLLS	120
ADI E112, S210	KPTHEMVEFMMSGITKYELGVESENELIVDPMPNLYFTRDPFASVGNVGVTHHEMRYIVRR	180
ADI E112, S210	RETLFARFVFRNHPKLVKTPWYYDPAMKMSIEGGDVFIYNNETLVVGVSERTDLDTTILL	240
ADI E112, S210	AKNIKANKEVEFKRIVAINVPKWNTNLMHLDTWLTMLDKNKFLYSPIANDVFKFWDYDLVN	300
ADI E112, S210	GGAEPQPQLNGLPLDKLLASIINKEPVLIPGGAGATEMEIARETNFDGTNYLAIRPGLV	360
ADI E112, S210	IGYDRNEKTNAALKAAGITVLPFHGNQLSLGMGNARCMSPLSRKDVKW	409

PHOE0063.ST25.txt
SEQUENCE LISTING

<110> Phoenix Pharmacologics, Inc.

<120> Novel Mutated Form of Arginine Deiminase

<130> PHOE-0063

<150> 09/564,559

<151> 2000-05-04

<160> 4

<170> PatentIn version 3.0

<210> 1

<211> 409

<212> PRT

<213> Mycoplasma hominis

<400> 1

```

Met Ser Val Phe Asp Ser Lys Phe Asn Gly Ile His Val Tyr Ser Glu
1          5          10          15
Ile Gly Glu Leu Glu Thr Val Leu Val His Glu Pro Gly Arg Glu Ile
          20          25          30
Asp Tyr Ile Thr Pro Ala Arg Leu Asp Glu Leu Leu Phe Ser Ala Ile
          35          40          45
Leu Glu Ser His Asp Ala Arg Lys Glu His Gln Ser Phe Val Lys Ile
          50          55          60
Met Lys Asp Arg Gly Ile Asn Val Val Glu Leu Thr Asp Leu Val Ala
65          70          75          80
Glu Thr Tyr Asp Leu Ala Ser Lys Ala Ala Lys Glu Glu Phe Ile Glu
          85          90          95
Thr Phe Leu Glu Glu Thr Val Pro Val Leu Thr Glu Ala Asn Lys Lys
          100         105         110
Ala Val Arg Ala Phe Leu Leu Ser Lys Pro Thr His Glu Met Val Glu
          115         120         125
Phe Met Met Ser Gly Ile Thr Lys Tyr Glu Leu Gly Val Glu Ser Glu
          130         135         140
Asn Glu Leu Ile Val Asp Pro Met Pro Asn Leu Tyr Phe Thr Arg Asp
145         150         155         160
Pro Phe Ala Ser Val Gly Asn Gly Val Thr Ile His Phe Met Arg Tyr
          165         170         175
Ile Val Arg Arg Arg Glu Thr Leu Phe Ala Arg Phe Val Phe Arg Asn
          180         185         190
His Pro Lys Leu Val Lys Thr Pro Trp Tyr Tyr Asp Pro Ala Met Lys
          195         200         205
Met Pro Ile Glu Gly Gly Asp Val Phe Ile Tyr Asn Asn Glu Thr Leu
210         215         220
Val Val Gly Val Ser Glu Arg Thr Asp Leu Asp Thr Ile Thr Leu Leu
225         230         235         240

```

PHOE0063.ST25.txt

Ala Lys Asn Ile Lys Ala Asn Lys Glu Val Glu Phe Lys Arg Ile Val
 245 250 255

Ala Ile Asn Val Pro Lys Trp Thr Asn Leu Met His Leu Asp Thr Trp
 260 265 270

Leu Thr Met Leu Asp Lys Asn Lys Phe Leu Tyr Ser Pro Ile Ala Asn
 275 280 285

Asp Val Phe Lys Phe Trp Asp Tyr Asp Leu Val Asn Gly Gly Ala Glu
 290 295 300

Pro Gln Pro Gln Leu Asn Gly Leu Pro Leu Asp Lys Leu Leu Ala Ser
 305 310 315 320

Ile Ile Asn Lys Glu Pro Val Leu Ile Pro Ile Gly Gly Ala Gly Ala
 325 330 335

Thr Glu Met Glu Ile Ala Arg Glu Thr Asn Phe Asp Gly Thr Asn Tyr
 340 345 350

Leu Ala Ile Lys Pro Gly Leu Val Ile Gly Tyr Asp Arg Asn Glu Lys
 355 360 365

Thr Asn Ala Ala Leu Lys Ala Ala Gly Ile Thr Val Leu Pro Phe His
 370 375 380

Gly Asn Gln Leu Ser Leu Gly Met Gly Asn Ala Arg Cys Met Ser Met
 385 390 395 400

Pro Leu Ser Arg Lys Asp Val Lys Trp
 405

<210> 2
 <211> 409
 <212> PRT
 <213> Mycoplasma hominis

<400> 2

Met Ser Val Phe Asp Ser Lys Phe Asn Gly Ile His Val Tyr Ser Glu
 1 5 10 15

Ile Gly Glu Leu Glu Thr Val Leu Val His Glu Pro Gly Arg Glu Ile
 20 25 30

Asp Tyr Ile Thr Pro Ala Arg Leu Asp Glu Leu Leu Phe Ser Ala Ile
 35 40 45

Leu Glu Ser His Asp Ala Arg Lys Glu His Gln Ser Phe Val Lys Ile
 50 55 60

Met Lys Asp Arg Gly Ile Asn Val Val Glu Leu Thr Asp Leu Val Ala
 65 70 75 80

Glu Thr Tyr Asp Leu Ala Ser Lys Ala Ala Lys Glu Glu Phe Ile Glu
 85 90 95

Thr Phe Leu Glu Glu Thr Val Pro Val Leu Thr Glu Ala Asn Pro Glu
 100 105 110

Ala Val Arg Ala Phe Leu Leu Ser Lys Pro Thr His Glu Met Val Glu
 115 120 125

Phe Met Met Ser Gly Ile Thr Lys Tyr Glu Leu Gly Val Glu Ser Glu
 130 135 140

PHOE0063.ST25.txt

Asn Glu Leu Ile Val Asp Pro Met Pro Asn Leu Tyr Phe Thr Arg Asp
 145 150 155 160
 Pro Phe Ala Ser Val Gly Asn Gly Val Thr Ile His Phe Met Arg Tyr
 165 170 175
 Ile Val Arg Arg Arg Glu Thr Leu Phe Ala Arg Phe Val Phe Arg Asn
 180 185 190
 His Pro Lys Leu Val Lys Thr Pro Trp Tyr Tyr Asp Pro Ala Met Lys
 195 200 205
 Met Pro Ile Glu Gly Gly Asp Val Phe Ile Tyr Asn Asn Glu Thr Leu
 210 215 220
 Val Val Gly Val Ser Glu Arg Thr Asp Leu Asp Thr Ile Thr Leu Leu
 225 230 235 240
 Ala Lys Asn Ile Lys Ala Asn Lys Glu Val Glu Phe Lys Arg Ile Val
 245 250 255
 Ala Ile Asn Val Pro Lys Trp Thr Asn Leu Met His Leu Asp Thr Trp
 260 265 270
 Leu Thr Met Leu Asp Lys Asn Lys Phe Leu Tyr Ser Pro Ile Ala Asn
 275 280 285
 Asp Val Phe Lys Phe Trp Asp Tyr Asp Leu Val Asn Gly Gly Ala Glu
 290 295 300
 Pro Gln Pro Gln Leu Asn Gly Leu Pro Leu Asp Lys Leu Leu Ala Ser
 305 310 315 320
 Ile Ile Asn Lys Glu Pro Val Leu Ile Pro Ile Gly Gly Ala Gly Ala
 325 330 335
 Thr Glu Met Glu Ile Ala Arg Glu Thr Asn Phe Asp Gly Thr Asn Tyr
 340 345 350
 Leu Ala Ile Lys Pro Gly Leu Val Ile Gly Tyr Asp Arg Asn Glu Lys
 355 360 365
 Thr Asn Ala Ala Leu Lys Ala Ala Gly Ile Thr Val Leu Pro Phe His
 370 375 380
 Gly Asn Gln Leu Ser Leu Gly Met Gly Asn Ala Arg Cys Met Ser Met
 385 390 395 400
 Pro Leu Ser Arg Lys Asp Val Lys Trp
 405

<210> 3
 <211> 409
 <212> PRT
 <213> Mycoplasma hominis

<400> 3

Met Ser Val Phe Asp Ser Lys Phe Asn Gly Ile His Val Tyr Ser Glu
 1 5 10 15
 Ile Gly Glu Leu Glu Thr Val Leu Val His Glu Pro Gly Arg Glu Ile
 20 25 30
 Asp Tyr Ile Thr Pro Ala Arg Leu Asp Glu Leu Leu Phe Ser Ala Ile
 35 40 45

PHOE0063.ST25.txt

Leu Glu Ser His Asp Ala Arg Lys Glu His Gln Ser Phe Val Lys Ile
 50 55 60
 Met Lys Asp Arg Gly Ile Asn Val Val Glu Leu Thr Asp Leu Val Ala
 65 70 75 80
 Glu Thr Tyr Asp Leu Ala Ser Lys Ala Ala Lys Glu Glu Phe Ile Glu
 85 90 95
 Thr Phe Leu Glu Glu Thr Val Pro Val Leu Thr Glu Ala Asn Lys Lys
 100 105 110
 Ala Val Arg Ala Phe Leu Leu Ser Lys Pro Thr His Glu Met Val Glu
 115 120 125
 Phe Met Met Ser Gly Ile Thr Lys Tyr Glu Leu Gly Val Glu Ser Glu
 130 135 140
 Asn Glu Leu Ile Val Asp Pro Met Pro Asn Leu Tyr Phe Thr Arg Asp
 145 150 155 160
 Pro Phe Ala Ser Val Gly Asn Gly Val Thr Ile His Phe Met Arg Tyr
 165 170 175
 Ile Val Arg Arg Arg Glu Thr Leu Phe Ala Arg Phe Val Phe Arg Asn
 180 185 190
 His Pro Lys Leu Val Lys Thr Pro Trp Tyr Tyr Asp Pro Ala Met Lys
 195 200 205
 Met Ser Ile Glu Gly Gly Asp Val Phe Ile Tyr Asn Asn Glu Thr Leu
 210 215 220
 Val Val Gly Val Ser Glu Arg Thr Asp Leu Asp Thr Ile Thr Leu Leu
 225 230 235 240
 Ala Lys Asn Ile Lys Ala Asn Lys Glu Val Glu Phe Lys Arg Ile Val
 245 250 255
 Ala Ile Asn Val Pro Lys Trp Thr Asn Leu Met His Leu Asp Thr Trp
 260 265 270
 Leu Thr Met Leu Asp Lys Asn Lys Phe Leu Tyr Ser Pro Ile Ala Asn
 275 280 285
 Asp Val Phe Lys Phe Trp Asp Tyr Asp Leu Val Asn Gly Gly Ala Glu
 290 295 300
 Pro Gln Pro Gln Leu Asn Gly Leu Pro Leu Asp Lys Leu Leu Ala Ser
 305 310 315 320
 Ile Ile Asn Lys Glu Pro Val Leu Ile Pro Ile Gly Gly Ala Gly Ala
 325 330 335
 Thr Glu Met Glu Ile Ala Arg Glu Thr Asn Phe Asp Gly Thr Asn Tyr
 340 345 350
 Leu Ala Ile Lys Pro Gly Leu Val Ile Gly Tyr Asp Arg Asn Glu Lys
 355 360 365
 Thr Asn Ala Ala Leu Lys Ala Ala Gly Ile Thr Val Leu Pro Phe His
 370 375 380
 Gly Asn Gln Leu Ser Leu Gly Met Gly Asn Ala Arg Cys Met Ser Met
 385 390 395 400
 Pro Leu Ser Arg Lys Asp Val Lys Trp

PHOE0063.ST25.txt

405

<210> 4
 <211> 409
 <212> PRT
 <213> Mycoplasma hominis

<400> 4

```

Met Ser Val Phe Asp Ser Lys Phe Asn Gly Ile His Val Tyr Ser Glu
1          5          10          15
Ile Gly Glu Leu Glu Thr Val Leu Val His Glu Pro Gly Arg Glu Ile
20          25          30
Asp Tyr Ile Thr Pro Ala Arg Leu Asp Glu Leu Leu Phe Ser Ala Ile
35          40          45
Leu Glu Ser His Asp Ala Arg Lys Glu His Gln Ser Phe Val Lys Ile
50          55          60
Met Lys Asp Arg Gly Ile Asn Val Val Glu Leu Thr Asp Leu Val Ala
65          70          75
Glu Thr Tyr Asp Leu Ala Ser Lys Ala Ala Lys Glu Glu Phe Ile Glu
85          90          95
Thr Phe Leu Glu Glu Thr Val Pro Val Leu Thr Glu Ala Asn Lys Glu
100         105         110
Ala Val Arg Ala Phe Leu Leu Ser Lys Pro Thr His Glu Met Val Glu
115         120         125
Phe Met Met Ser Gly Ile Thr Lys Tyr Glu Leu Gly Val Glu Ser Glu
130         135         140
Asn Glu Leu Ile Val Asp Pro Met Pro Asn Leu Tyr Phe Thr Arg Asp
145         150         155
Pro Phe Ala Ser Val Gly Asn Gly Val Thr Ile His Phe Met Arg Tyr
165         170         175
Ile Val Arg Arg Arg Glu Thr Leu Phe Ala Arg Phe Val Phe Arg Asn
180         185         190
His Pro Lys Leu Val Lys Thr Pro Trp Tyr Tyr Asp Pro Ala Met Lys
195         200         205
Met Ser Ile Glu Gly Gly Asp Val Phe Ile Tyr Asn Asn Glu Thr Leu
210         215         220
Val Val Gly Val Ser Glu Arg Thr Asp Leu Asp Thr Ile Thr Leu Leu
225         230         235
Ala Lys Asn Ile Lys Ala Asn Lys Glu Val Glu Phe Lys Arg Ile Val
245         250         255
Ala Ile Asn Val Pro Lys Trp Thr Asn Leu Met His Leu Asp Thr Trp
260         265         270
Leu Thr Met Leu Asp Lys Asn Lys Phe Leu Tyr Ser Pro Ile Ala Asn
275         280         285
Asp Val Phe Lys Phe Trp Asp Tyr Asp Leu Val Asn Gly Gly Ala Glu
290         295         300
Pro Gln Pro Gln Leu Asn Gly Leu Pro Leu Asp Lys Leu Leu Ala Ser

```

Page 5

305 310 PHOE0063.ST25.txt 315 320
Ile Ile Asn Lys Glu Pro Val Leu Ile Pro Ile Gly Gly Ala Gly Ala
 325 330 335
Thr Glu Met Glu Ile Ala Arg Glu Thr Asn Phe Asp Gly Thr Asn Tyr
 340 345 350
Leu Ala Ile Lys Pro Gly Leu Val Ile Gly Tyr Asp Arg Asn Glu Lys
 355 360 365
Thr Asn Ala Ala Leu Lys Ala Ala Gly Ile Thr Val Leu Pro Phe His
 370 375 380
Gly Asn Gln Leu Ser Leu Gly Met Gly Asn Ala Arg Cys Met Ser Met
385 390 395 400
Pro Leu Ser Arg Lys Asp Val Lys Trp
 405

**This Page is Inserted by IFW Indexing and Scanning
Operations and is not part of the Official Record**

BEST AVAILABLE IMAGES

Defective images within this document are accurate representations of the original documents submitted by the applicant.

Defects in the images include but are not limited to the items checked:

☒ **BLACK BORDERS**

☐ **IMAGE CUT OFF AT TOP, BOTTOM OR SIDES**

☐ **FADED TEXT OR DRAWING**

☒ **BLURRED OR ILLEGIBLE TEXT OR DRAWING**

☐ **SKEWED/SLANTED IMAGES**

☐ **COLOR OR BLACK AND WHITE PHOTOGRAPHS**

☐ **GRAY SCALE DOCUMENTS**

☐ **LINES OR MARKS ON ORIGINAL DOCUMENT**

☐ **REFERENCE(S) OR EXHIBIT(S) SUBMITTED ARE POOR QUALITY**

☐ **OTHER:** _____

IMAGES ARE BEST AVAILABLE COPY.

As rescanning these documents will not correct the image problems checked, please do not report these problems to the IFW Image Problem Mailbox.